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SEQUENCE LISTING

<110> Rowe, Peter

<120> A Novel Polypeptide Hormone Phosphatonin

<130> BEAR-008

<140> US 09/700,696

<141> 2000-11-17

<150> PCT EP99/03403

<151> 1999-05-18

<150> GB 9810681.8

<151> 1998-05-18

<150> GB 9819387.3

<151> 1998-09-04

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| ctacatgacc | aagaagaata | tggcgcaagct | ctcatcagaa | ataacatgc | acatataatg | 180 |
| gggcaggatga | ctgcgattaa | actcctgggg | gaagaaaaca | aagagaacac | accttaggaat | 240 |
| gttctaaaca | taatcccagc | aagtatgaat | tatgctaaag | cacactcgaa | ggataaaaag | 300 |
| aagcctcaaa | gagattccca | agcccgaaaa | agtccagtaa | aaagcaaaag | caccatcg | 360 |
| attcaacaca | acattgacta | cctaaaacat | ctctcaaaag | tcaaaaaaat | ccccagtgtat | 420 |
| tttgaaggca | gcgggttatac | agatcttcaa | gagagagggg | acaatgat | atctccttgc | 480 |
| agtggggacg | gcacaccttt | taaggacatt | cctgttaaag | gagaagctac | tggcctgac | 540 |
| ctagaaggca | aagatattca | aacagggttt | gcaggccaa | gtgaagctga | gagtaactcat | 600 |
| cttgacaccaa | aaaagccagg | ttataatgag | atcccaagaga | gagaagaaaa | tggtgaaat | 660 |
| accattggaa | ctagggatga | aactgcgaaa | gaggcagatg | ctgttgatgt | cagccttgta | 720 |
| gagggcagca | acgatatcat | gggtgttacc | aattttaaagg | agctccctgg | aagagaagga | 780 |
| aacagagttg | atgctggcag | ccaaaatgt | caccaaggga | aggttgagtt | tcattaccct | 840 |
| cctgcaccct | caaaaagagaa | aagaaaagaa | ggcagtagtg | atgcagctga | aagtaccaac | 900 |
| tataatgaaa | ttcctaaaaaa | tggcaaaggc | agtaccagaa | agggtgtaga | tcattctaat | 960 |
| aggaaccaag | caacctaaaa | tgaaaaacaa | agtttctta | gtaaggccaa | aagtcaaggc | 1020 |
| ctgcccattc | cttctcggtg | tcttgataat | gaaatcaaaa | acgaaatgg | ttccttaat | 1080 |
| ggccccagtc | atgagaatat | aataacacat | ggcagaaaaat | atcattatgt | accccacaga | 1140 |
| caaaaataatt | ctacacggaa | taagggtatg | ccacaaggga | aaggctctg | ggtagacaa | 1200 |
| ccccattcca | acaggagggtt | tagtcccgt | agaagggatg | acagtagtg | gtcatctgac | 1260 |
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| agtctgaaga | cctcgacc | tgtgagttga | tgttagaggag | agccacctga | cagctgacca | 1380 |
| ggtgaagaga | ggataagagt | aagaactgag | tgagccaaga | atcctggct | ccttggggga | 1440 |
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35 40 45
Ala Ala Leu Ile Arg Asn Asn Met Gln His Ile Met Gly Pro Val Thr
50 55 60
Ala Ile Lys Leu Leu Gly Glu Asn Lys Glu Asn Thr Pro Arg Asn
65 70 75 80
Val Leu Asn Ile Ile Pro Ala Ser Met Asn Tyr Ala Lys Ala His Ser
85 90 95
Lys Asp Lys Lys Pro Gln Arg Asp Ser Gln Ala Gln Lys Ser Pro
100 105 110
Val Lys Ser Lys Ser Thr His Arg Ile Gln His Asn Ile Asp Tyr Leu
115 120 125
Lys His Leu Ser Lys Val Lys Ile Pro Ser Asp Phe Glu Gly Ser
130 135 140
Gly Tyr Thr Asp Leu Gln Glu Arg Gly Asp Asn Asp Ile Ser Pro Phe
145 150 155 160
Ser Gly Asp Gly Gln Pro Phe Lys Asp Ile Pro Gly Lys Gly Glu Ala
165 170 175
Thr Gly Pro Asp Leu Glu Gly Lys Asp Ile Gln Thr Gly Phe Ala Gly
180 185 190
Pro Ser Glu Ala Glu Ser Thr His Leu Asp Thr Lys Lys Pro Gly Tyr
195 200 205
Asn Glu Ile Pro Glu Arg Glu Glu Asn Gly Gly Asn Thr Ile Gly Thr
210 215 220
Arg Asp Glu Thr Ala Lys Glu Ala Asp Ala Val Asp Val Ser Leu Val
225 230 235 240
Glu Gly Ser Asn Asp Ile Met Gly Ser Thr Asn Phe Lys Glu Leu Pro
245 250 255
Gly Arg Glu Gly Asn Arg Val Asp Ala Gly Ser Gln Asn Ala His Gln
260 265 270
Gly Lys Val Glu Phe His Tyr Pro Pro Ala Pro Ser Lys Glu Lys Arg
275 280 285
Lys Glu Gly Ser Ser Asp Ala Ala Glu Ser Thr Asn Tyr Asn Glu Ile
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Pro Lys Asn Gly Lys Gly Ser Thr Arg Lys Gly Val Asp His Ser Asn
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340 345 350
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Thr His Gly Arg Lys Tyr His Tyr Val Pro His Arg Gln Asn Asn Ser
370 375 380

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